

- TNTN

C →

D4-1 →

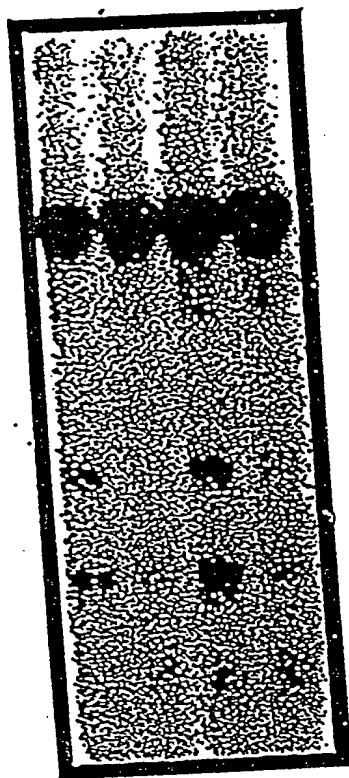


FIGURE 1

1 cttcaaagtt ccccgcatga aaattactta aacgttgcac acaacgttcc gaaaaatott
 61 ttgtgaaga agaaaaggaa attcagtgtg tgagtctcag caggagttaa gtaattgcag
 121 ctttaataa tccogaaaaa gaagcgctta tctgcgggca gagtgccttc gattctcttc
 181 ctgtgccaga tgattagtgc actggaagta cctcttgatc caaaacttct tgaagacttg
 241 gtacagcctc caaccatcac ccaacagtot ccaaaagatt acattattga cctcgggag
 301 aatattgtaa tccagtgtga agccaaaggg aaaccgcccc caagcttttc ctggaccgt
 361 aatgggactc attttgacat cgataaagac cctctggtca ccatgaagcc tggcacagga
 421 acgtcctataa ttaacatcat gagcgaaggg aaagotgaga cctatgaagg agtctatcag
 481 tgtacagcaa ggaacgaacg cggagctgca gtttctaata acattgttgt ccgcccaccc
 541 agatcaccat tgtggaccaa agaaaaactt gaaccaatca cacttcaaag tggcagtot
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 661 aattcctttc aaagacttcc acaaagttag agagtttttc aaggtttgaa tggggacott
 721 tatttttcca atgtcctccc agaggacacc cgogaagact atatctgtta tctagattt
 781 aatcatactc aaaccataca gcagaagcaa cctatttctg tgaaggtgat ttcagtggat
 841 gaattgaatg acactatagc tgctaatttg agtgacactg agttttatgg tgctaaatca
 901 agtagagaga ggccaccaac atttttaact ccagaaggca atgcaagtaa caaagaggaa
 961 ttaagaggaa atgtgctttc actggagtgc attgcagaag gactgcctac ccaattatt
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 1141 aaaaatgcat taggagccat ccactatacc atttctgtta gagttaaagc ggtccatcac
 1201 tggatcacag cccctcaaaa tcttgtgctg tcccaggag aggatgggag cttgatctgc
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 1381 caagaaagat caagtgcagt atatcagtg tatgcctcta atgaattatg atattactg
 1441 gcaaacgcat ttgtaaatgt gctggctgag ccaaccagaa tctcacacc tgcaaacaca
 1501 ctctaccagg tcattgcaaa caggcctgct ttactagact gtgccttctt tgggtctcot
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 2821 atcctcacct tccaaggcag caagactcat ggcattgttc cggggctaga gcccttttagc
 2881 cactacacac tgaatgtccg agtgggtcaat gggaaagggg agggcccagc cagccctgac
 2941 agagtcttta atactccaga aggagtcctc agtgotccct cgtctttgaa gattgtgaat
 3001 ccaacactgg actctctcac tttggaatgg gatccaccga gccaccogaa tggcattttg
 3061 acagagtaca ccttaaagta tcagocaaat aacagocac atgaattagg ccccttggta
 3121 gatttgaaaa ttcctgccaa caagacacgg tggactttaa aaaatttaa tttcagcact
 3181 cgtatataag tttattttota tgcaaaaaca tcagcaggat caggaaagta aattacagag
 3241 gaagcagtaa caactgtgga tgaagctggt attcttocac ctgatgtagg tgcaggcaaa
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 3481 ttaaagggtc taatgccagg aacagcctac aaagtctgag ttggtgctgt gggggactot
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 3661 attttgtga ttgtttgctt cctcagaaga aacaagggtg gtaaatatcc agttaaagaa
 3721 aaggaagatg cccatgctga cctgaaatc cagcctatga aggaagatga tgggacattt
 3781 ggagaataca gtgatgcaga agaccacaag cctttgaaa aaggaagtcg aactccttca
 3841 gacaggactg tgaaaaaaga agatagtac gacagocag ttgactatgt agaaggggtt
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 4021 tttgtttat ttttaagctc aaagccaata ttgatttct ctagaatgtt tatcctaagc
 4081 tcttgtttgt cagccctctc atagtatga atgtgggtg gagagtatat tttc

FIGURE 2A

BT110
BT181

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 QOSPKEYIIDPRENIVIQCEAKGKPPPSFSWTRNGTHFDIDKDPLVTMKPGTGTLIIN
 IMSEGKAETIEGVYQCTARNERGAAVSNNIVVRPSRSPILWTKLEPITLQSGQSLVL
 PCRPPIGLPPPIIFWMDNSFORLPQSERVSQGLNGDLYFSNVLPEDTREDYICYARFN
 HTQTIQKQKQISVKVISVDELNDTIAANLSDTEFYGAKSSRERPPTFLTPEGNASNKE
 ELRGNVLSLECIAEGLPTPIIYWAKEDGMLPKNRTVYKNFEKTLQIIHVSEADSGNYQ
 CIAKNALGAIHHTISVRVKAAPYWITAPQNLVLS PGEDGT LICRANGNPKPRISWLTN
 GVPIEIAPDPSRKIDGDTIIFSNVQERS SAVYQCNASNEYGYLLANAFVNVLAEPFR
 ILTPANTLYQVIANRPALDCAFFGSPLPTIEWFKGAKGSALHEDIYVLHENGTLK
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 DHTLSLTVLWLKDNRELPSEDERETVDKDLVADVSDDDSGTYTCVANTTLDVSVASA
 VLSVVAPTPTPAFVYDVNPFPDLELTDQLDKSVQLSWTPGDDNNSPITKFIIEYEDA
 MHKPGWLWHHQTEVSGTQTTAQLKLSFYVNYSEFVMAVNSIGKSLPSEASEQYLTASE
 PDKNPTAVEGLGSEPDNLEITWKPLNGFESNGPGLQYKVSQRQKDGDEWTSVVVANV
 SKYIVSGTPTFVPLYLIKVQALNDMGFAPEPAVVMGHSGEDLPMVAPGNVRVNVNSTL
 AEVHWDVPVPLKSIRGHLQGYRIYYWKTQSSSKRNRRIEKKILTFOGSKTHGMLPGL
 PFSHYTLNVRVVGKGEPPASPDVRVNTPEGVPSAPSSLKIVNPTLDSLTLEWDPSSH
 PNGILTEYTLKYQPINSTHELGLVLDLKI PANKTRWTLKNLNFSTRYKFFYQTSAG
 SGSQITEEA VTTVDEAGILPPDVGAGKVQAVNTRISNLTA AAAETIANISWEYEGPEH
 VNEYVEYGVAGSKEWRKEIVNGSRSEFGLKGLMPGTAYKVRVGAVGDSGFVSSDVF
 ETGPAMASRQVDIATQGWFIGIMCAVALLILILLIVCFIRRNKGGKYPVKEKEDAHAD
 PEIQPMKEDDGTGEYS DAEDHKPLKKSRTPSDRTVKKEDSDDSLVDYGEVNGQFN
 EDGSFIGQYSGKKEKEPAEGNESSEAPSFVNAMNSFV"

FIGURE 2B

HUMAN Nr-CAM	4097	TCTCATACTATGAACATATGGGTAGAGAGTATATTTTC	4134


D4-1	1	TCTCATACTATGAACATATGGGTAGAGAGTATATTTTCTGCTGT	44
		***** ** ** ***** **	
RAT Nr-CAM	3713	TCTCATACTATGGACATATGGGTAGAAAGAATGTTTTCTGOGGT	3757
D4-1	45	ATGTTAGTATTATGAGAATAGTTACAGCAAAACATAACTCAGT	89
		** * ***** **** ** ** * * * * *	
RAT Nr-CAM	3758	ATATGAGTATTATAAGAACAGAGCAAGAACATAACTCAGTCAGT	3802
D4-1	90	CAAAGTATATGTTAATATGAACTGGAATGCAAAGTGCATACTT	134
		** * *** ***** ** *** *	
RAT Nr-CAM	3802	CAGATGATACGTTAATATGAACTGGGGTGAAAAGG	3831
D4-1	135	TTTCATTCAAATGGGTATTCTTGATTTCCTAAAAAAAAAAAAA	179


FIGURE 2C


hNr-CAM

hNr-CAM1/3AS



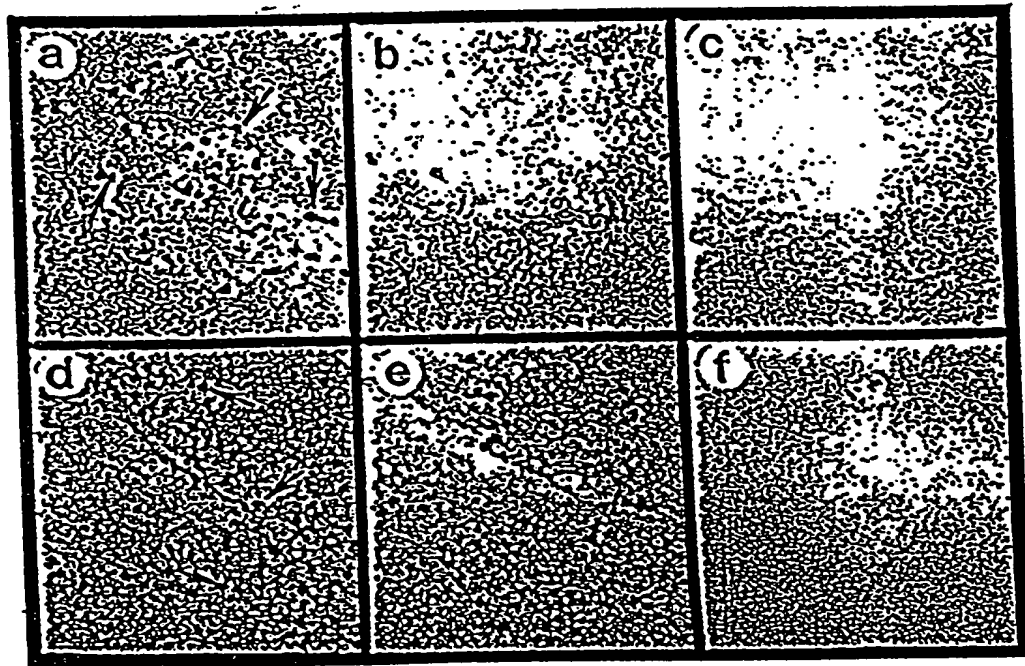
 Ig domain

 Fibronectin domain

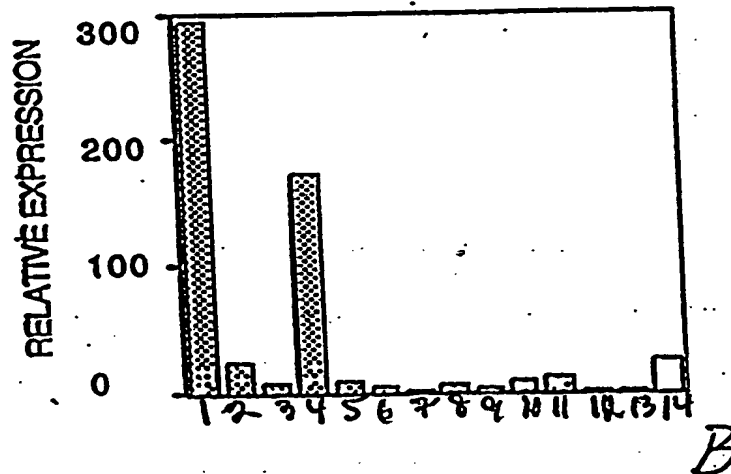
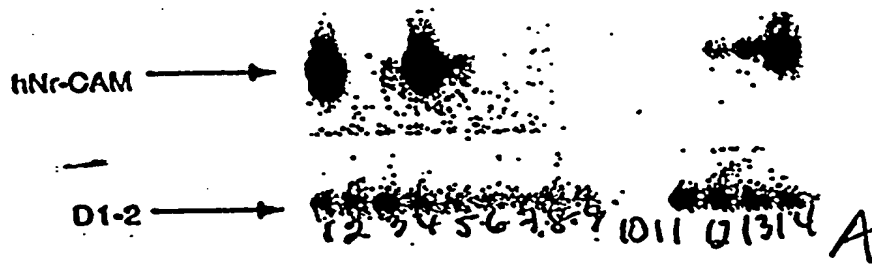
 Transmembrane domain

 Cytoplasmic tail

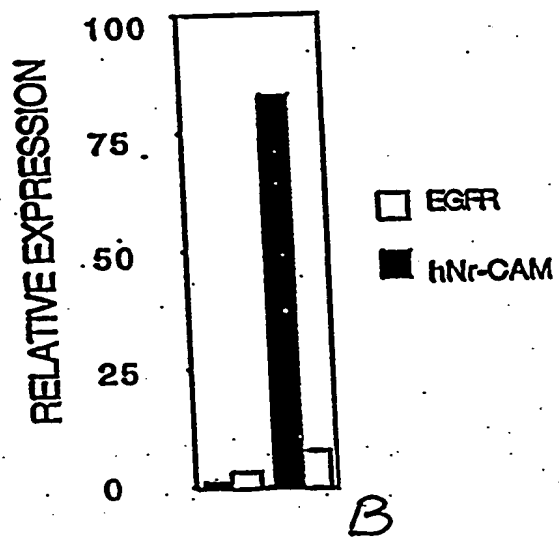
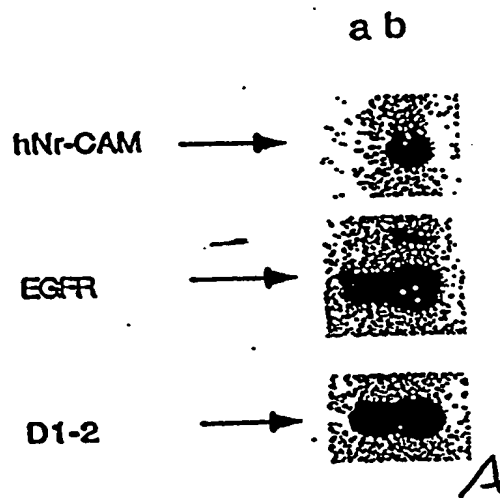
FIGURE 2D



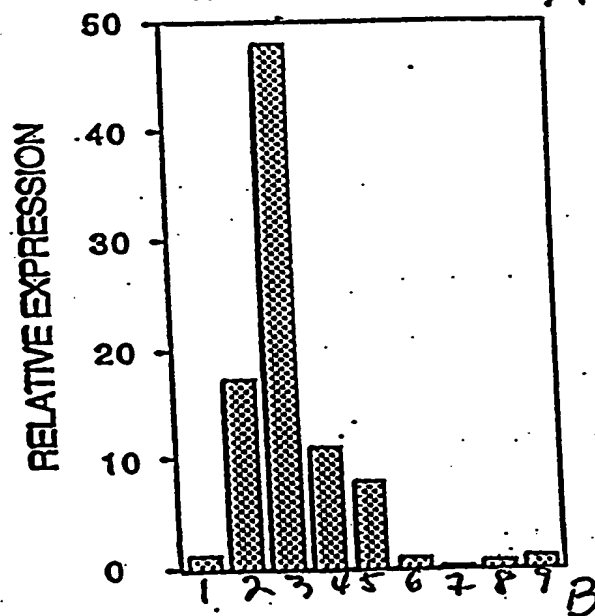
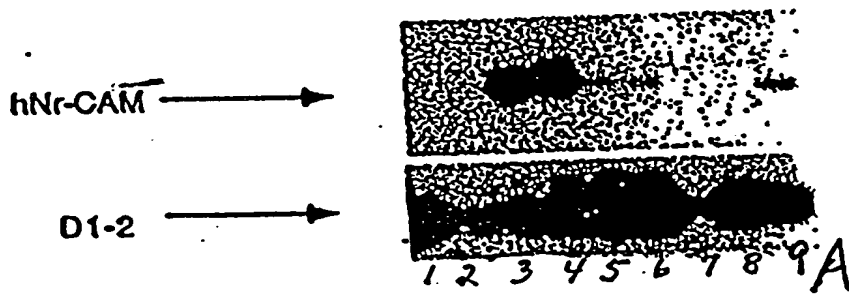
FIGURES 3A-F



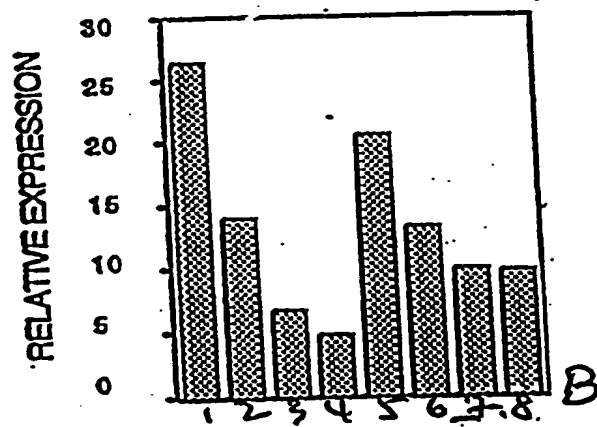
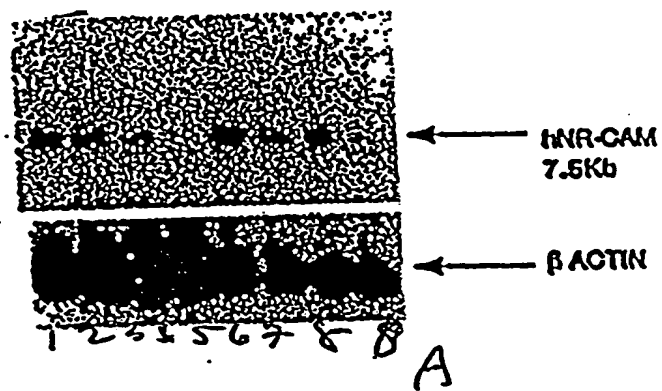
FIGURES 4A-B



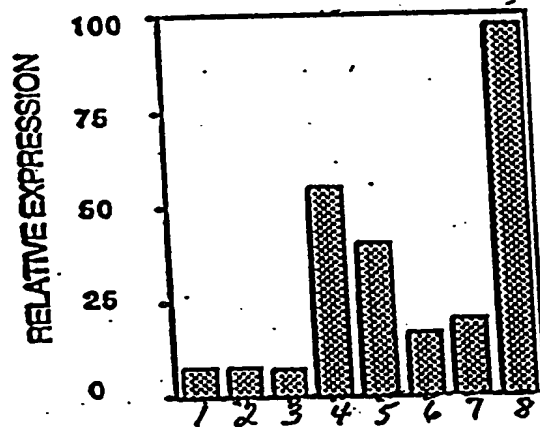
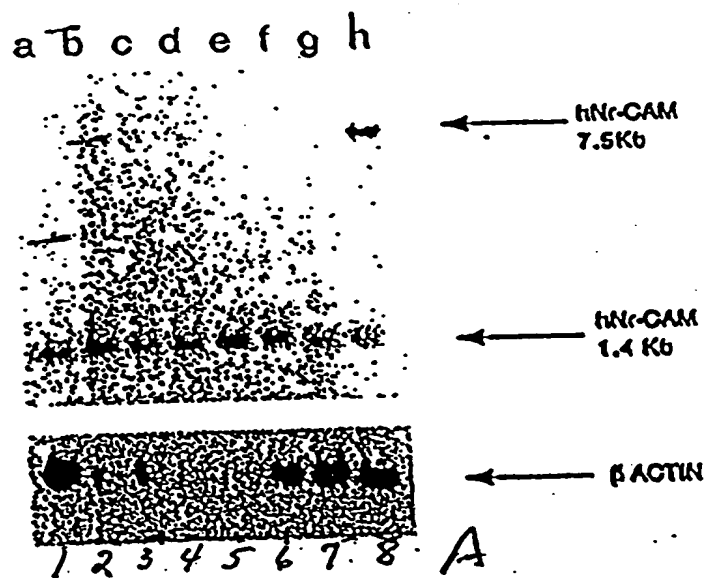
FIGURES 5A-B



FIGURES 6A-B



FIGURES 7A-B



FIGURES 8A-B

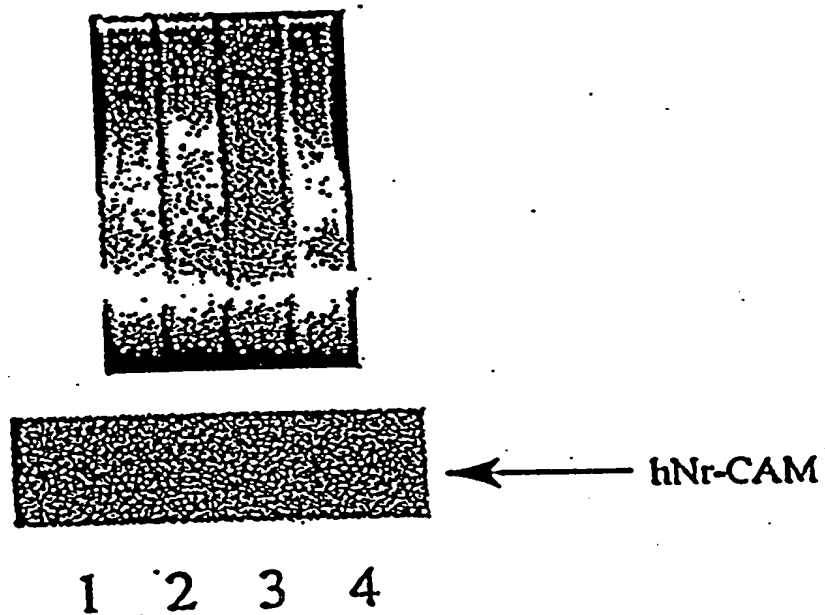
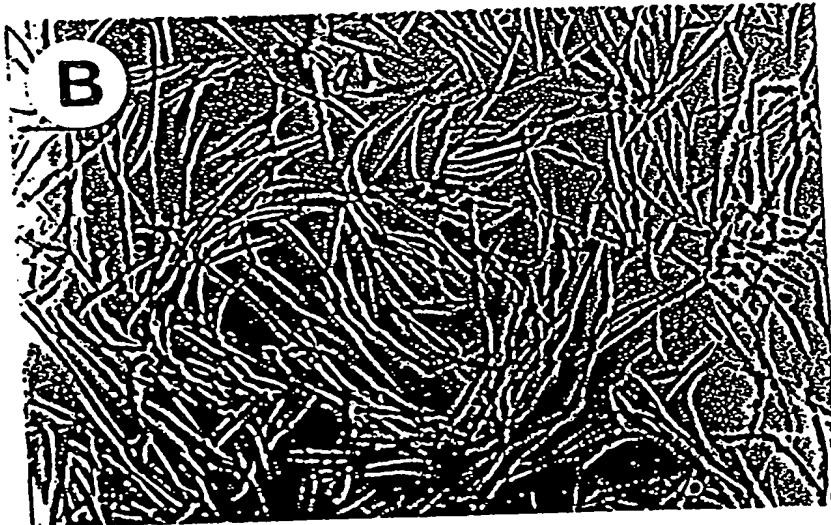
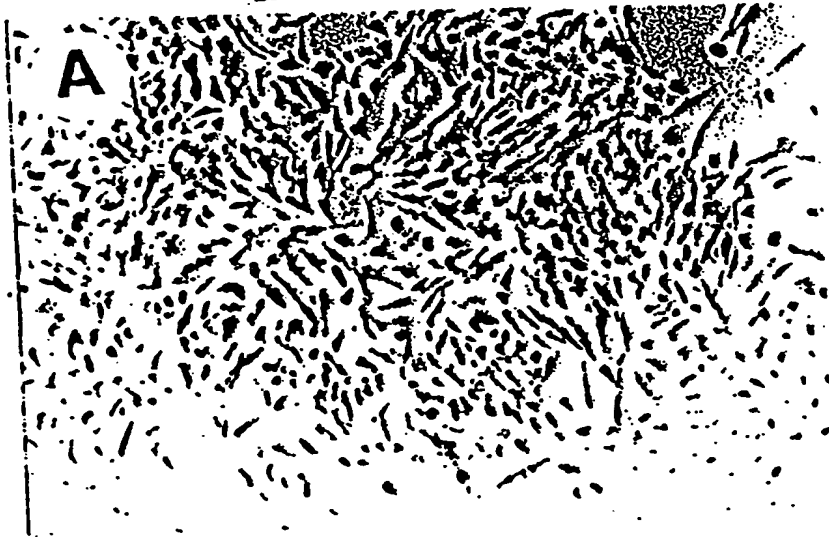


FIGURE 9



FIGURES 10A-B

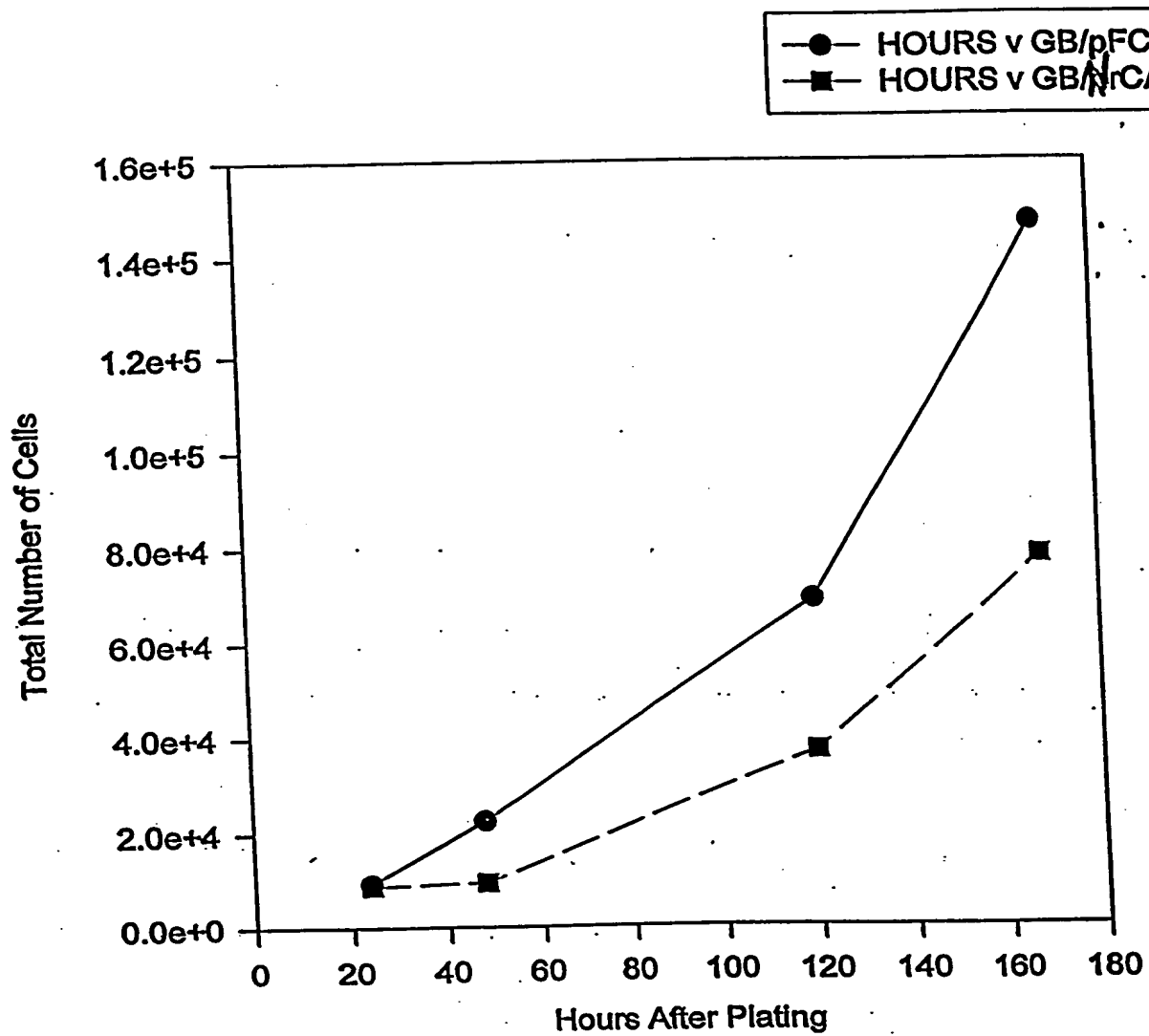


FIGURE 11

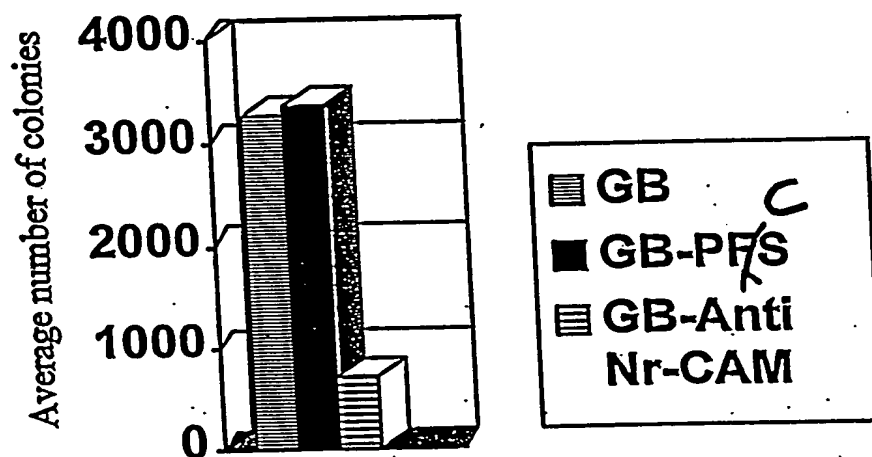


FIGURE 12

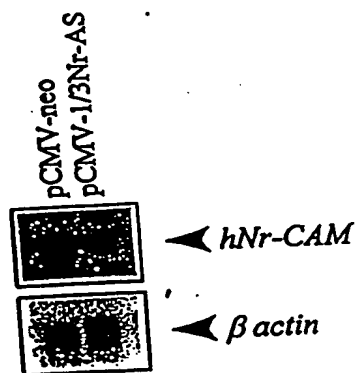


FIGURE 13

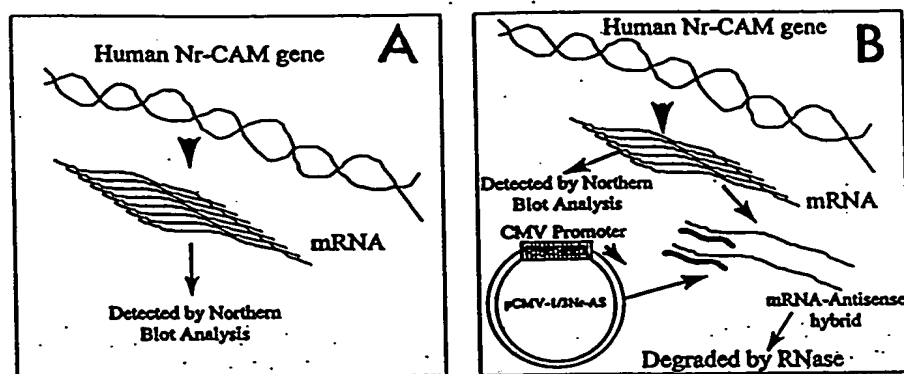


FIGURE 14A-B

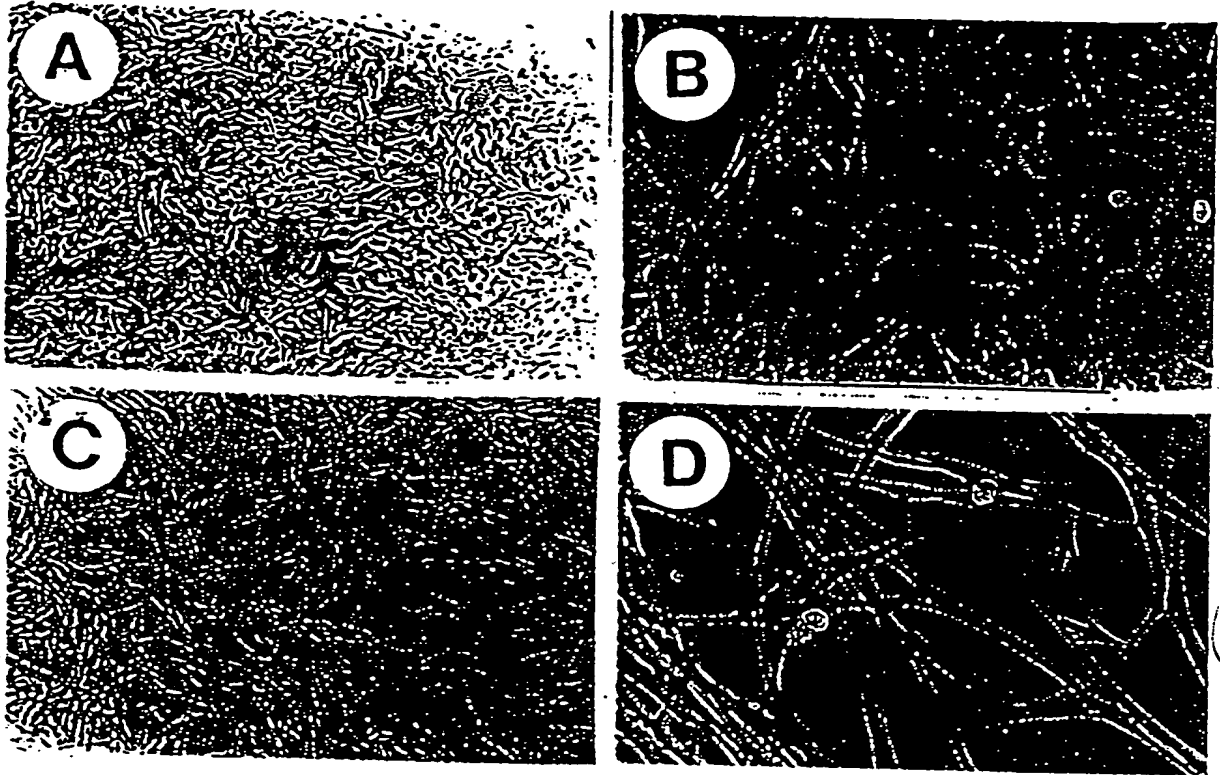


FIGURE 15A-D

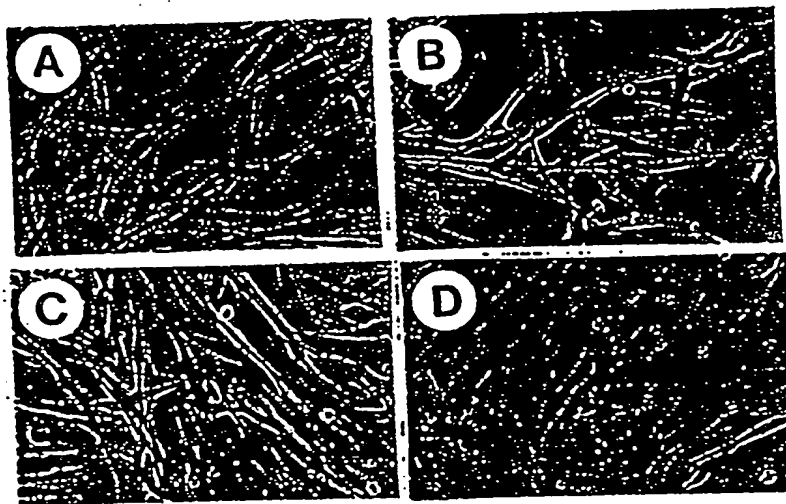


FIGURE 16A-D

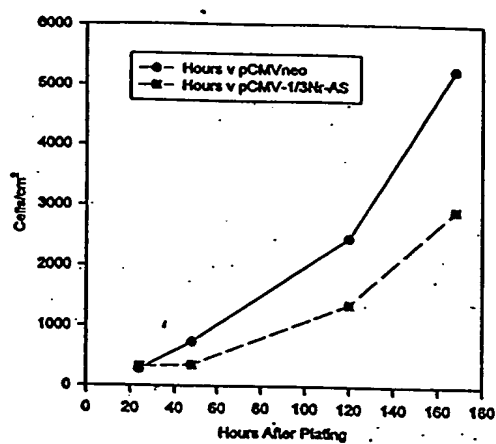


FIGURE 17

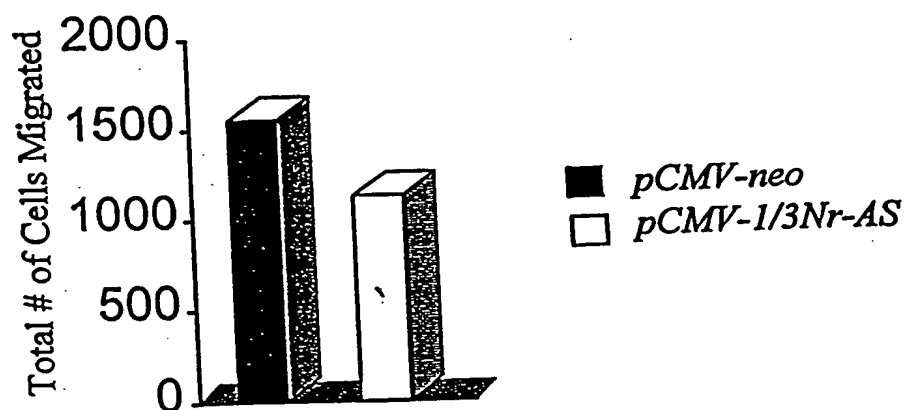


FIGURE 18

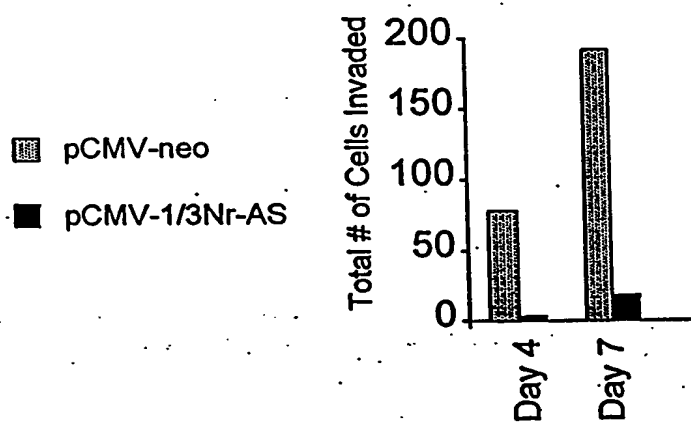


FIGURE 19

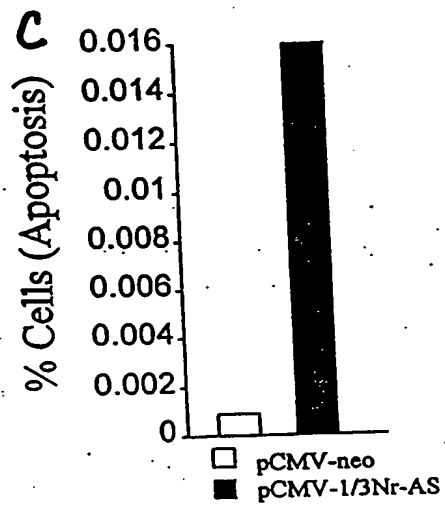
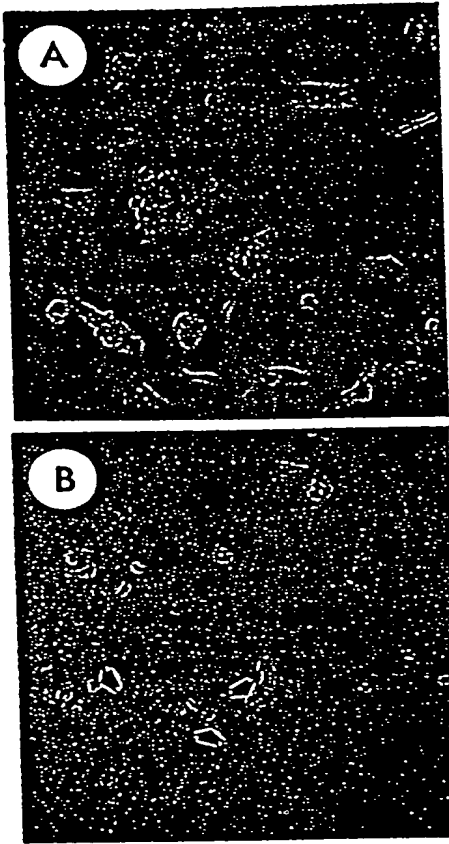


FIGURE 20A-C



FIGURE 21

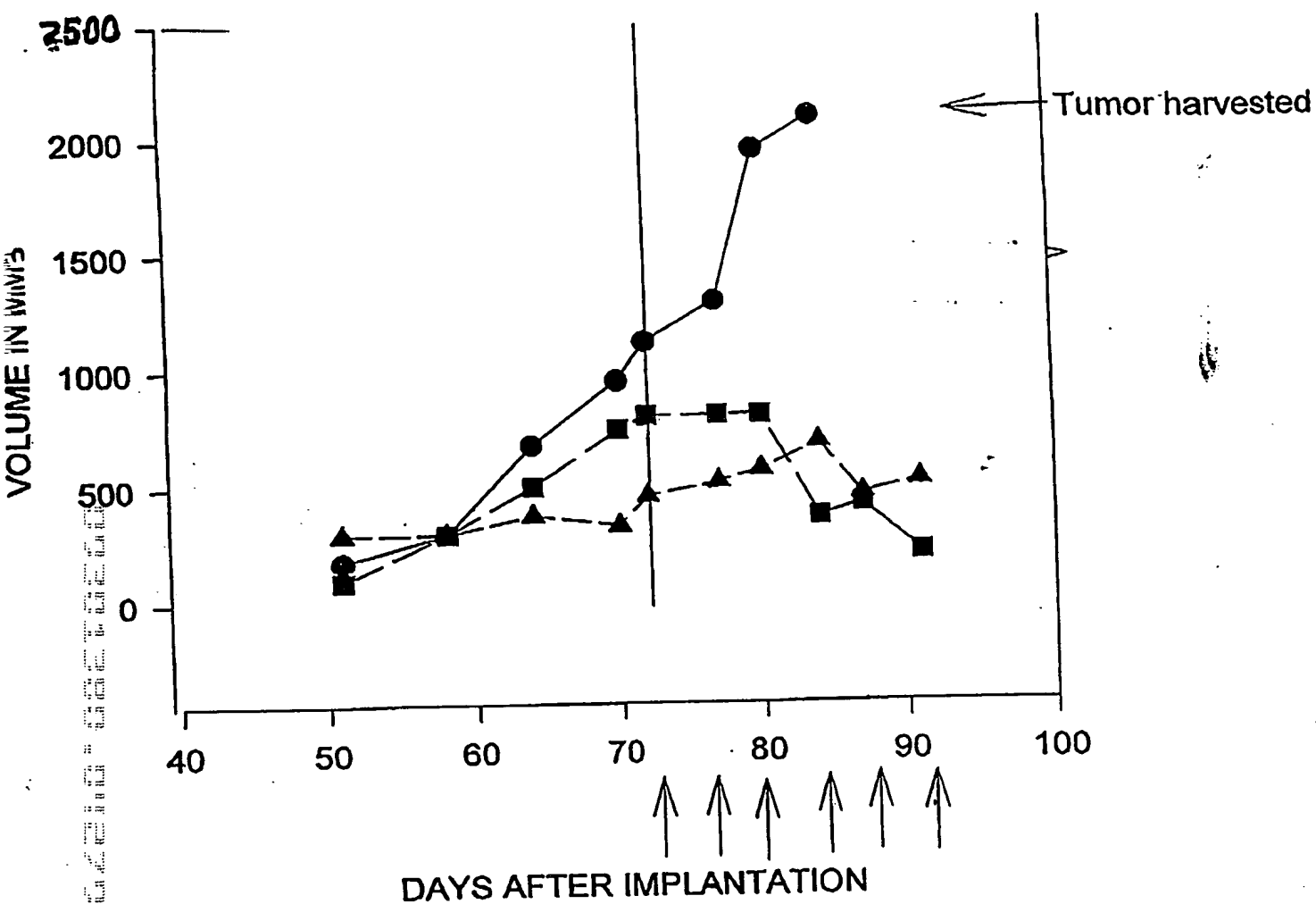


FIGURE 22A

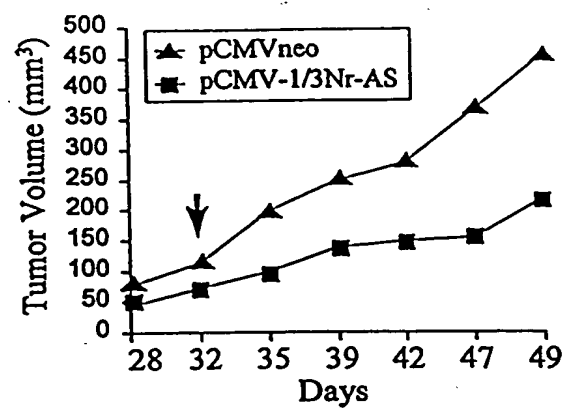
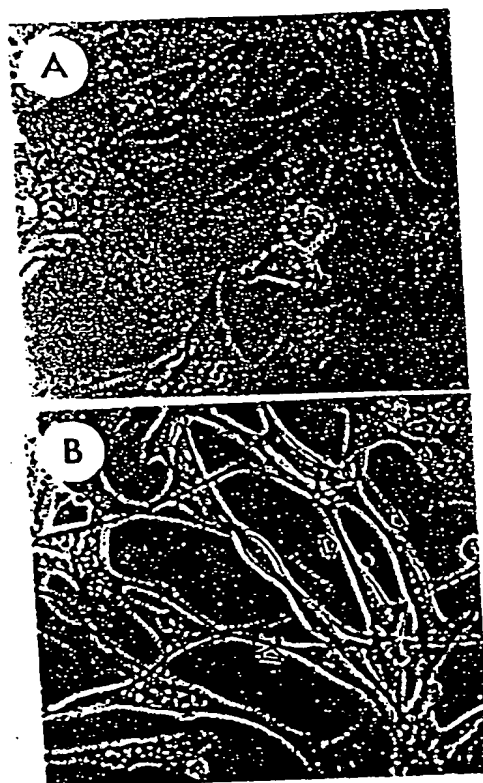


FIGURE 22B



Average # of colonies (1×10^5)

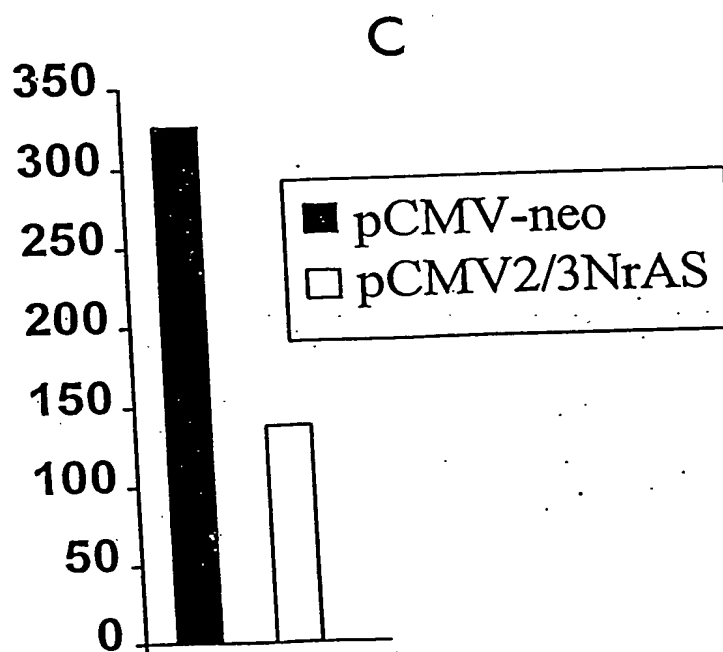


FIGURE 23A-C

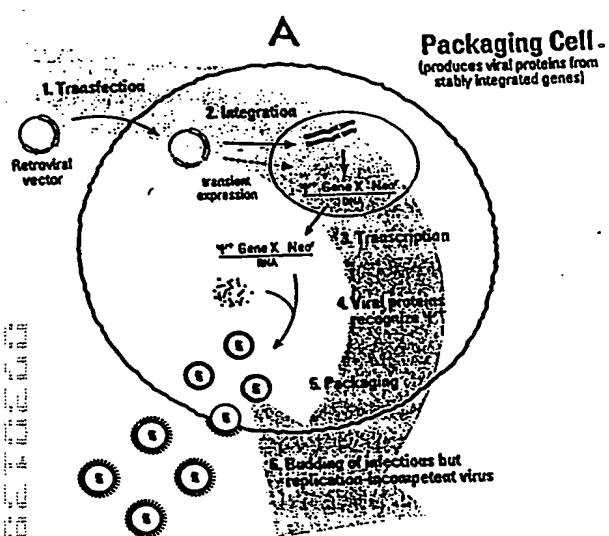


FIGURE 25A

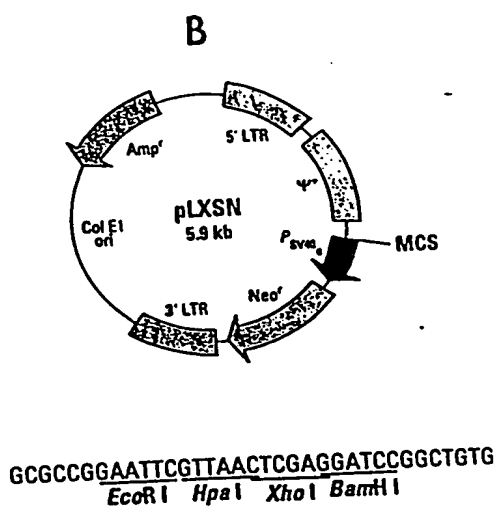


FIGURE 25B

pCMV-1/3Nr-AS

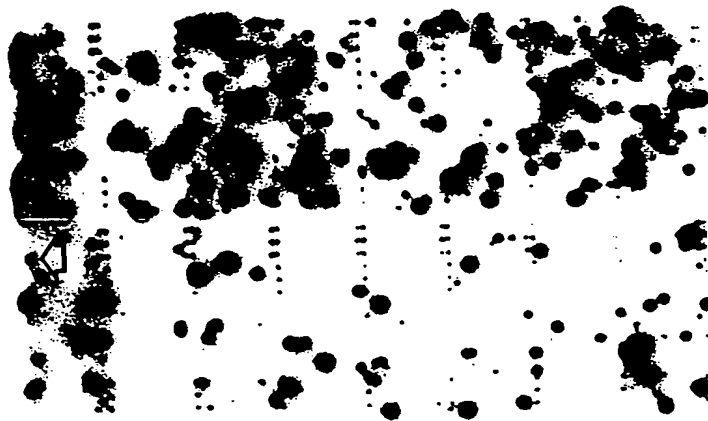


FIGURE 26A

pCMV-neo

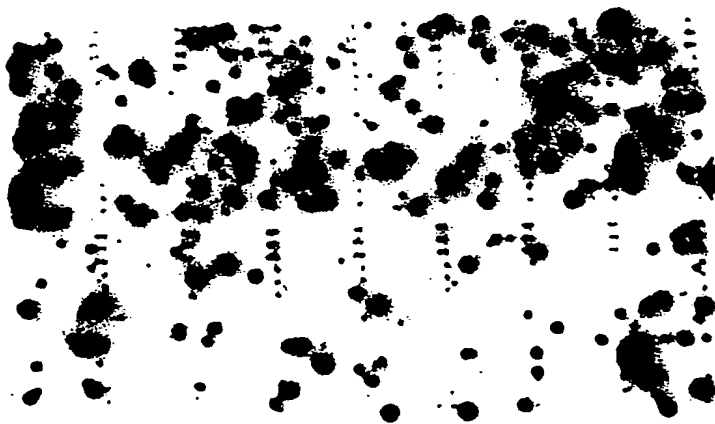


FIGURE 26B